

Resource Summary Report

Generated by NIF on Sep 4, 2026

PRINTS

RRID:SCR_003412

Type: Tool

Proper Citation

PRINTS (RRID:SCR_003412)

Resource Information

URL: <http://www.bioinf.man.ac.uk/dbbrowser/PRINTS/>

Proper Citation: PRINTS (RRID:SCR_003412)

Description: Compendium of protein fingerprints. Diagnostic fingerprint database.

Abbreviations: PRINTS

Resource Type: data or information resource, database

Keywords: compedium, protein, fingerprint, diagnostic, database, FASEB list

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: PRINTS

Resource ID: SCR_003412

Alternate IDs: nif-0000-03338

Alternate URLs: <http://130.88.97.239/PRINTS/index.php>

Record Creation Time: 20220129T080218+0000

Record Last Update: 20260904T000117+0000

Ratings and Alerts

No rating or validation information has been found for PRINTS.

No alerts have been found for PRINTS.

Data and Source Information

Usage and Citation Metrics

We found 427 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Zhang L, et al. (2026) Quercetin upregulates steroid hormone biosynthesis to enhance reproductive performance in roosters. *Poultry science*, 105(4), 106590.

Chen J, et al. (2026) Phylogenomics-driven metalloantibiotic engineering: Overcoming ciprofloxacin resistance in *Pseudomonasaeruginosa* with sideromycin-bismuth synergy. *Cell reports. Medicine*, 7(3), 102637.

Jiang K, et al. (2026) Telomere-to-telomere-level genome assembly and annotation of the Zi goose *Anser cygnoides*. *Scientific data*, 13(1).

Du Z, et al. (2026) High-resolution phage-host assignment through key proteins using large language models. *Nature communications*, 17(1).

Ouyang Y, et al. (2026) Chromosome-scale Genome Assembly of the Critically Endangered Blue-crowned Laughingthrush (*Pterorhinus courtoisi*, Leiothrichidae). *Scientific data*, 13(1).

Pan Y, et al. (2026) Proteomic Analysis of Lotus-Derived NnAP2 Regulation of Soluble Sugar and Starch Content in Potato (*Solanum tuberosum*). *Plants (Basel, Switzerland)*, 15(4).

Soman SP, et al. (2026) IsoProDB: an integrated map of human protein isoforms for accelerated research. *Database : the journal of biological databases and curation*, 2026.

Lu D, et al. (2026) An annotated genome of freshwater amphipod (*Gammarus nekkensis*). *Scientific data*, 13(1).

K??stlbacher S, et al. (2026) Prediction of eukaryotic cellular complexity in Asgard archaea using structural modelling. *Nature microbiology*, 11(3), 747.

Liu T, et al. (2026) Chromosome-level genome assembly and annotation of the endangered plant *Primula kwangtungensis* (Primulaceae). *Scientific data*, 13(1), 212.

Yu ZX, et al. (2026) Chromosome-level genome of *Mosla chinensis* from alpine ecotype provides insights into terpenoid biosynthesis and germplasm exploration. *Frontiers in plant science*, 17, 1819008.

Zou R, et al. (2026) A chromosome-level genome assembly of *Cistanche deserticola* provides insights into its evolution and molecular mechanisms of parasitism. *Plant communications*, 7(1), 101581.

Huang Z, et al. (2026) Integrated Proteomics and Metabolomics Analysis Reveals Potential Pathways Underlying Onion-Mediated Regulation of Spleen Immune Function in

Liangshan Black Sheep. Veterinary sciences, 13(5).

Liber JA, et al. (2026) Aimea gen. nov. defines a novel plant-associated yeast genus in Microbotryomycetes with three novel species. bioRxiv : the preprint server for biology.

Yin S, et al. (2026) A telomere-to-telomere genome assembly of *Castanopsis orthacantha* (Fagaceae). Scientific data, 13(1).

Nneji LM, et al. (2026) First Chromosome-level Genome Assembly and Annotation of an Endangered Freshwater Stingray (*Fontitrygon garouaensis*) from Africa. Scientific data, 13(1).

De Panis D, et al. (2026) The reference genome of the Asian Elephant (*Elephas maximus*): a foundation for conservation and genomic research. BMC genomics, 27(1).

Stilwell PA, et al. (2026) The genome of *Istocheta aldrichi* (Diptera: Tachinidae), a parasitoid of the Japanese beetle, *Popillia japonica* (Coleoptera: Scarabaeidae). G3 (Bethesda, Md.), 16(4).

Sedaghatkish A, et al. (2026) A contamination-reduced genome of *Plasmodiophora brassicae* with comprehensive functional annotation. BMC microbiology, 26(1), 118.

Dos Santos Menezes F, et al. (2026) Comparative analysis of PRR receptors and R genes reveals structural and functional divergence between wild, ancestral, and commercial citrus species. BMC genomics, 27(1), 173.